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RAW SEQUENCE LISTING DATE: 03/12/2002 PATENT APPLICATION: US/10/084,206 TIME: 14:02:32

Input Set : A:\191D1C1.ST25.txt

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3 <110> APPLICANT: Li et al.
     5 <120> TITLE OF INVENTION: G-Protein Receptor HTNAD29
     7 <130> FILE REFERENCE: PF191D1C1
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/084,206
C--> 9 <141> CURRENT FILING DATE: 2002-02-28
     9 <150> PRIOR APPLICATION NUMBER: US95/07288
    10 <151> PRIOR FILING DATE: 1995-06-06
    12 <150> PRIOR APPLICATION NUMBER: 08/468,534
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    15 <150> PRIOR APPLICATION NUMBER: 09/399,095
    16 <151> PRIOR FILING DATE: 1999-09-20
    18 <160> NUMBER OF SEQ ID NOS: 9
    20 <170> SOFTWARE: PatentIn Version 3.1
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    23 <211> LENGTH: 1753
    24 <212> TYPE: DNA
    25 <213> ORGANISM: Homo sapiens
    27 <220> FEATURE:
    28 <221> NAME/KEY: CDS
    29 <222> LOCATION: (523)..(1533)
    30 <223> OTHER INFORMATION:
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    42 ataatgaaca catggacttc tgtttattaa attcagttga cccctttagc caattgccag
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    46 tttaaaagca ttgatgaatt aattagaact ttagacaaca agaaaaattg aaaagaattc
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                                                                              480
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    50 ctttctaaga actaatataa ttgctacctt aaaaaggaaa aa atg aac agc aca
                                                                              534
    51
                                                       Met Asn Ser Thr
    52
                                                                              582
    54 tgt att gaa gaa cag cat gac ctg gat cac tat ttg ttt ccc att gtt
    55 Cys Ile Glu Glu Gln His Asp Leu Asp His Tyr Leu Phe Pro Ile Val
                            10
                                                15
    58 tac atc ttt gtg att ata gtc agc att cca gcc aat att gga tct ctg
                                                                              630
    59 Tyr Ile Phe Val Ile Ile Val Ser Ile Pro Ala Asn Ile Gly Ser Leu
    60
                        25
                                            30
                                                                              678
    62 tgt gtg tct ttc ctg caa ccc aag aag gaa agt gaa cta gga att tac
    63 Cys Val Ser Phe Leu Gln Pro Lys Lys Glu Ser Glu Leu Gly Ile Tyr
    64
                    40
                                        45
    66 ctc ttc agt ttg tca cta tca gat tta ctc tat gca tta act ctc cct.
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Input Set : A:\191D1C1.ST25.txt

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71 Leu Trp Ile Asp Tyr Thr Trp Asn Lys Asp Asn Trp Thr Phe Ser Pro 72 70 75 80	
74 gcc ttg tgc aaa ggg agt gct ttt ctc atg tac atg aag ttt tac agc	822
75 Ala Leu Cys Lys Gly Ser Ala Phe Leu Met Tyr Met Lys Phe Tyr Ser	022
76 85 90 95 100	
78 age aca gea tte etc ace tge att gee gtt gat egg tat ttg get gtt	870
79 Ser Thr Ala Phe Leu Thr Cys Ile Ala Val Asp Arg Tyr Leu Ala Val	0.0
80 105 110 115	
82 gtc tac cct ttg aag ttt ttt ttc cta agg aca aga aga att gca ctc	918
83 Val Tyr Pro Leu Lys Phe Phe Leu Arg Thr Arg Arg Ile Ala Leu	
84 120 125 130	
86 atg gtc agc ctg tcc atc tgg ata ttg gaa acc atc ttc aat gct gtc	966
87 Met Val Ser Leu Ser Ile Trp Ile Leu Glu Thr Ile Phe Asn Ala Val	
88 135 140 145	
90 atg ttg tgg gaa gat gaa aca gtt gtt gaa tat tgc gat gcc gaa aag	1014
91 Met Leu Trp Glu Asp Glu Thr Val Val Glu Tyr Cys Asp Ala Glu Lys	
92 150 155 160	
94 tct aat ttt act tta tgc tat gac aaa tac cct tta gag aaa tgg caa	1062
95 Ser Asn Phe Thr Leu Cys Tyr Asp Lys Tyr Pro Leu Glu Lys Trp Gln	
96 165 170 175 180	
98 atc aac ctc aac ttg ttc agg acg tgt aca ggc tat gca ata cct ttg	1110
99 Ile Asn Leu Asn Leu Phe Arg Thr Cys Thr Gly Tyr Ala Ile Pro Leu	
100 185 190 195	
102 gtc acc atc ctg atc tgt aac cgg aaa gtc tac caa gct gtg cgg cac	1158
103 Val Thr Ile Leu Ile Cys Asn Arg Lys Val Tyr Gln Ala Val Arg His	
104 200 205 210	
106 aat aaa gcc acg gaa aac aag gaa aag aag aga atc ata aaa cta ctt	1206
107 Asn Lys Ala Thr Glu Asn Lys Glu Lys Lys Arg Ile Ile Lys Leu Leu	
108 215 220 225	1054
110 gtc agc atc aca gtt act ttt gtc tta tgc ttt act ccc ttt cat gtg	1254
111 Val Ser Ile Thr Val Thr Phe Val Leu Cys Phe Thr Pro Phe His Val	
112 230 235 240	1 202
114 atg ttg ctg att cgc tgc att tta gag cat gct gtg aac ttc gaa gac	1302
115 Met Leu Leu Ile Arg Cys Ile Leu Glu His Ala Val Asn Phe Glu Asp 116 245 250 255 260	
116 245 250 255 260 118 cac agc aat tet ggg aag cga act tac aca atg tat aga atc acg gtt	1350
119 His Ser Asn Ser Gly Lys Arg Thr Tyr Thr Met Tyr Arg Ile Thr Val	1330
120 265 270 275	
122 gca tta aca agt tta aat tgt gtt gct gat cca att ctg tac tgt ttt	1398
123 Ala Leu Thr Ser Leu Asn Cys Val Ala Asp Pro Ile Leu Tyr Cys Phe	1370
124 280 285 290	
126 gtt acc gaa aca gga aga tat gat atg tgg aat ata tta aaa ttc tgc	1446
127 Val Thr Glu Thr Gly Arg Tyr Asp Met Trp Asn Ile Leu Lys Phe Cys	_110
128 295 300 305	
130 act ggg agg tgt aat aca tca caa aga caa aga aaa cgc ata ctt tct	1494
131 Thr Gly Arg Cys Asn Thr Ser Gln Arg Gln Arg Lys Arg Ile Leu Ser	

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Input Set : A:\191D1C1.ST25.txt

132		310					315					320					
134	gtg	tct	aca	aaa	gat	act	atg	gaa	tta	gag	gtc	ctt	gag	taga	aacca	aag	1543
135	Val	Ser	Thr	Lys	Asp	Thr	Met	Glu	Leu	Glu	Val	Leu	Glu				
136	6 325 330 335																
138	gate	gttt	tga a	aggga	aagg	ga ag	gttta	aagti	t ate	gcat	tatt	ata	tcate	caa	gatta	acattt	1603
140	tgaa	aaag	gaa a	atcta	agcat	tg t	gaggg	ggact	gact aagtgttctc			agagtgatgt tttaatccag					1663
	42 tocaataaaa atatottaaa actgoattgt acagotoco												1723				
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155					5					10					15		
158	Phe	Pro	Ile	Val	Tyr	Ile	Phe	Val	Ile	Ile	Val	Ser	Ile	Pro	Ala	Asn	
159				20					25					30			
162	Ile	Gly	Ser	Leu	Cys	Val	Ser	Phe	Leu	Gln	Pro	Lys	Lys	Glu	Ser	Glu	
163		-	35		_			40				_	45				
166	Leu	Gly	Ile	Tyr	Leu	Phe	Ser	Leu	Ser	Leu	Ser	Asp	Leu	Leu	Tyr	Ala	
167		50					55					60			_		
170	Leu	Thr	Leu	Pro	Leu	Trp	Ile	Asp	Tyr	Thr	Trp	Asn	Lys	Asp	Asn	Trp	
171						70					75					80	
174	Thr	Phe	Ser	Pro	Ala	Leu	Cys	Lys	Gly	Ser	Ala	Phe	Leu	Met	Tyr	Met	
175					85					90					95		
178	Lys	Phe	Tyr	Ser	Ser	Thr	Ala	Phe	Leu	Thr	Cys	Ile	Ala	Val	Asp	Arg	
179				100					105					110			
182	Tyr	Leu	Ala	Val	Val	Tyr	Pro	Leu	Lys	Phe	Phe	Phe	Leu	Arg	Thr	Arg	
183			115					120					125				
186	Arg	Ile	Ala	Leu	Met	Val	Ser	Leu	Ser	Ile	Trp	Ile	Leu	Glu	Thr	Ile	
187		130					135					140					
190	Phe	Asn	Ala	Val	Met	Leu	Trp	Glu	Asp	Glu	Thr	Val	Val	Glu	Tyr	Cys	
191	145					150					155					160	
194	Asp	Ala	Glu	Lys	Ser	Asn	Phe	Thr	Leu	Cys	Tyr	Asp	Lys	Tyr	Pro	Leu	
195					165					170					175		
198	Glu	Lys	Trp	Gln	Ile	Asn	Leu	Asn	Leu	Phe	Arg	Thr	Cys	Thr	Gly	Tyr	
199				180					185					190			
202	Ala	Ile	Pro	Leu	Val	Thr	Ile	Leu	Ile	Cys	Asn	Arg	Lys	Val	Tyr	Gln	
203			195					200					205				
206	Ala	Val	Arg	His	Asn	Lys	Ala	Thr	Glu	Asn	Lys	Glu	Lys	Lys	Arg	Ile	
207		210					215					220					
210	Ile	Lys	Leu	Leu	Val	Ser	Ile	Thr	Val	Thr	Phe	Val	Leu	Cys	Phe	Thr	
211	225					230					235					240	
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Input Set : A:\191D1C1.ST25.txt

Output Set: N:\CRF3\03122002\J084206.raw

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Input Set : A:\191D1C1.ST25.txt

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322
            290
                                 295
                                                     300
    325 Tyr Ser Met Arg Ser Ser Arg Lys Cys Ser Arg Ala Thr Thr Asp Thr
    326 305
                            310
                                                315
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                         325
    330
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    341 <400> SEQUENCE: 4
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    342 cgaattcctc catgaacagc acatgtatt
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    376
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    378 <400> SEQUENCE: 7
                                                                                60
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                                                                                61
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    393 <400> SEQUENCE: 8
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    397 <210> SEQ ID NO: 9
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VERIFICATION SUMMARY DATE: 03/12/2002 PATENT APPLICATION: US/10/084,206 TIME: 14:02:34

Input Set : A:\191D1C1.ST25.txt

Output Set: N:\CRF3\03122002\J084206.raw

 $\hbox{L:9 M:270 C: Current Application Number differs, Replaced Current Application No}\\$

L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date